

**AMENDMENTS TO THE CLAIMS:**

This listing of claims replaces all prior versions and listings of claims in the application:

1-14. (Canceled).

15. (Currently Amended) A method of characterising a treatment applied to a population of cells, comprising:

providing a treatment to a population of cells;

deriving a plurality of cellular features from at least a first captured image of the population of cells that have been exposed to the treatment;

creating an on-target effect signature, which is characteristic of an on-target effect of the treatment on the population of cells, from ~~at least a first one of~~ a group of cellular features selected from the plurality of cellular features derived from the population of cells, the ~~at least one of the plurality of~~ group of cellular features relating to cellular properties involved in the on-target effect;

creating a side effect signature, which is characteristic of a side effect to the on-target effect, from ~~at least a second one of~~ a second group of cellular features selected from the plurality of cellular features derived from the population of cells, the second ~~one of the plurality~~ group of cellular features relating to cellular properties not being involved in the on-target effect;

creating an on-target effect metric derived from the on-target effect signature;  
creating a side effect metric derived from the side effect signature; and  
comparing the on-target effect metric to the side effect metric to thereby  
characterise the response of the population of cells to the treatment.

16. (Canceled).

17. (Currently Amended) The method as claimed in claim ~~[[16]]~~15, wherein ~~the side effect signature is created from a further group of cellular features, in which~~  
none of the members of the group of cellular features used to create the on-target effect  
signature and the members of the further group of cellular features used to created the  
side effect signature are common.

18. (Currently Amended) The method as claimed in claim 15, wherein the  
second ~~one~~ group of the ~~plurality~~ of cellular features is affected by the treatment.

19. (Canceled)

20. (Previously Presented) The method as claimed in claim 15, wherein  
deriving one or both of the on-target effect metric and the side effect metric includes  
determining the difference between the on-target effect signature or side effect

signature and a control signature from the same cellular features for a control group of cells.

21. (Original) The method as claimed in claim 15, and further comprising:  
capturing at least a first image of a control group of cells; and  
deriving a plurality of cellular features from the image of the control group of cells;  
creating a control on-target signature for the same cellular features for the control group; and  
creating a control side effect signature for the same cellular features for the control group.

22. (Original) The method of claim 21, further comprising determining a side effect distance in a multivariate space between the side effect signature and the control side effect signature.

23. (Previously Presented) The method of claim 22, further comprising determining an on-target effect distance in a multivariate space between the on-target effect signature and the control on-target effect signature.

24. (Previously Presented) The method of claim 23, wherein characterising the treatment is based on the side effect distance.

25. (Previously Presented) The method of claim 24, wherein characterising the treatment is also based on the on-target effect distance.

26. (Original) The method as claimed in claim 25, further comprising generating a graphical representation of the side effect distance and on-target effect distance.